## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/583,59/
Source:	IFWP.
Date Processed by STIC:	6/27/06

## ENTERED



DATE: 06/27/2006

**IFWP** 

PATENT APPLICATION: US/10/583,591 TIME: 10:43:38 Input Set : A:\PA0394 Sequence Listing.ST25.txt Output Set: N:\CRF4\06272006\J583591.raw 3 <110> APPLICANT: STUBBS, Simon L. FRANCIS, Michael J. CUSHING, Adrian 5 ISMAIL, Rahman A. 8 <120> TITLE OF INVENTION: CYTOCHROME C PROTEIN AND ASSAY 10 <130> FILE REFERENCE: PA0394 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/583,591 13 <141> CURRENT FILING DATE: 2006-06-19 15 <150> PRIOR APPLICATION NUMBER: PCT/GB2004/005317 16 <151> FRIOR FILING DATE: 2004-12-17 18 <150> PRIOR APPLICATION NUMBER: GB 0329353.7 19 <151> PRIOR FILING DATE: 2003-12-19 21 <160> NUMBER OF SEQ ID NOS: 15 23 <170> SOFTWARE: PatentIn version 3.3 25 <210> SEO ID NO: 1 26 <211> LENGTH: 315 27 <212> TYPE: DNA 28 <213> ORGANISM: Homo sapiens 30 <400> SEQUENCE: 1 31 atgggtgatg ttgagaaagg caagaagatt tttattatga agtgttccca gtgccacacc 60 33 gttgaaaagg gaggcaagca caagactggg ccaaatctcc atggtctctt tgggcggaag 120 35 acaggtcagg cccctggata ctcttacaca gccgccaata agaacaaagg catcatctgg 180 37 ggagaggata cactgatgga gtatttggag aatcccaaga agtacatccc tggaacaaaa 39 atgatetttg teggeattaa gaagaaggaa gaaagggeag aettaatage ttateteaaa 300 315 41 aaagctacta atgag 44 <210> SEQ ID NO: 2 45 <211> LENGTH: 105 46 <212> TYPE: PRT 47 <213> ORGANISM: Homo sapiens 49 <400> SEQUENCE: 2 51 Met Gly Asp Val Glu Lys Gly Lys Lys Ile Phe Ile Met Lys Cys Ser 10 55 Gln Cys His Thr Val Glu Lys Gly Gly Lys His Lys Thr Gly Pro Asn 20 25 59 Leu His Gly Leu Phe Gly Arg Lys Thr Gly Gln Ala Pro Gly Tyr Ser 40 63 Tyr Thr Ala Ala Asn Lys Asn Lys Gly Ile Ile Trp Gly Glu Asp Thr 67 Leu Met Glu Tyr Leu Glu Asn Pro Lys Lys Tyr Ile Pro Gly Thr Lys 75 71 Met Ile Phe Val Gly Ile Lys Lys Glu Glu Arg Ala Asp Leu Ile 90

RAW SEQUENCE LISTING

75 Ala Tyr Leu Lys Lys Ala Thr Asn Glu

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Input Set : A:\PA0394 Sequence Listing.ST25.txt

Output Set: N:\CRF4\06272006\J583591.raw

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92 aaacttaccc ttaaatttat ttgcactact ggaaaactac ctgttccatg gccaacactt
94 gtcactactc tctcttatgg tgttcaatgc ttttcaagat acccagatca tatgaaacgg
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118 atcatctggg gagaggatac actgatggag tatttggaga atcccgccaa gtacatccct
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151 Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg
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159 Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val
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163 Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile
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PATENT APPLICATION: US/10/583,591 TIME: 10:43:38

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Output Set: N:\CRF4\06272006\J583591.raw

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175 Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val
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                                        170
179 Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro
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                                    185
183 Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser
184
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187 Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Gly Phe Val
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191 Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys Leu Glu
192 225
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                                             235
195 Asn Ser Thr Met Gly Asp Val Glu Lys Gly Lys Lys Ile Phe Ile Met
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199 Lys Cys Ser Gln Cys His Thr Val Glu Lys Gly Gly Lys His Lys Thr
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                                280
                                                     285
207 Gly Tyr Ser Tyr Thr Ala Ala Asn Lys Asn Lys Gly Ile Ile Trp Gly
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211 Glu Asp Thr Leu Met Glu Tyr Leu Glu Asn Pro Ala Lys Tyr Ile Pro
212 305
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242 aaagctacta atgagggtcg accegggatg agtaaaggag aagaactttt cactggagtt
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244 gtcccaattc ttgttgaatt agatggtgat gttaatgggc acaaattttc tgtcagtgga
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RAW SEQUENCE LISTING DATE: 06/27/2006
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323       Val Gln Cys       Phe Ser Arg       Tyr       Pro Asp His Met       Lys       Arg His Asp Phe         324       180       180       185       190       190         327       Phe Lys       Ser Ala Met       Pro Glu Gly Tyr       Val Gln Glu Arg Thr Ile Phe         328       195       200       205       205         331       Phe Lys       Asp Asp Gly Asn Tyr Lys       Thr Arg Ala Glu Val Lys Phe Glu         332       210       215       220         335       Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys         336       225       230       235         339       Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser         340       245       250         245       250       250         240       255         343       His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val         344       260       265         250       270         347       Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala         348       275       280         280       285         351       Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu	319	Lys	Leu	Pro	Val	Pro	Trp	Pro	Thr	Leu	Val	Thr	Thr	Leu	Ser	Tyr	Gly	
324       180       185       190         327 Phe Lys       Ser Ala Met       Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe         328       195       200       205         331 Phe Lys       Asp Asp Gly Asn Tyr Lys       Thr Arg Ala Glu Val Lys Phe Glu         332       210       215       220         335 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys       240         336 225       230       235       240         339 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser       240         340       245       250       255         343 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val       255         344 Lys Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala       265         347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Fro Val Leu Leu	320					165					170					175		
327       Phe Lys       Ser       Ala Met       Pro       Glu       Gly       Tyr       Val       Glu       Arg       Thr       Ile       Phe         328       195       200       200       205       205       205         331       Phe Lys       Asp       Asp       Gly       Asn       Tyr       Lys       Thr       Arg       Ala Glu       Val       Lys       Phe       Glu         332       210       215       215       220       220       220       220       220       220       220       220       220       230       220       235       240       240       240       235       240       240       235       240       240       240       235       240       240       240       235       240<	323	Val	Gln	Cys	Phe	Ser	Arg	Tyr	Pro	Asp	His	Met	Lys	Arg	His	Asp	Phe	
328       195       200       205         331 Phe Lys Asp Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu       220         332 210       215       220         335 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys       235         336 225       230       235         339 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser       240         340       245       250         250       255         343 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val         344 Lys Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala         347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala         348 275       280       285         351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu	324				180					185					190			
331 Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu 332 210 215 220  335 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys 336 225 230 235 240  339 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser 340 245 250 255  343 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val 344 260 265 270  347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala 348 275 280 285  351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu	327	Phe	Lys	Ser	Ala	Met	Pro	Glu	Gly	Tyr	Val	Gln	Glu	Arg	Thr	Ile	Phe	
332 210 215 220  335 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys 336 225 230 235 240  339 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser 340 245 250 255  343 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val 344 260 265 270  347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala 348 275 280 285  351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu	328			195					200					205				
332 210 215 220  335 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys 336 225 230 235 240  339 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser 340 245 250 255  343 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val 344 260 265 270  347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala 348 275 280 285  351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu	331	Phe	Lys	Asp	Asp	Gly	Asn	Tyr	Lys	Thr	Arg	Ala	Glu	Val	Lys	Phe	Glu	
336       225       230       235       240         339       Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser         340       245       250       255         343       His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val         344       260       265         347       Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala         348       275       280         351       Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu			_	_	_	-		-	-		_				_			
336       225       230       235       240         339       Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser         340       245       250       255         343       His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val         344       260       265         347       Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala         348       275       280         351       Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu	335	Gly	Asp	Thr	Leu	Val	Asn	Arq	Ile	Glu	Leu	Lys	Gly	Ile	Asp	Phe	Lys	
339 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser 340			_					_							_			
340       245       250       255         343 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val       260       265       270         347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala       280       285         351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu	339	Glu	Asp	Gly	Asn	Ile		Gly	His	Lys	Leu	Glu	Tyr	Asn	Tyr	Asn	Ser	
343 His Asn Val       Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val         344			-	-				-		•			•		-			
344 260 265 270  347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala  348 275 280 285  351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu		His	Asn	Val	Tyr		Met	Ala	Asp	Lys		Lys	Asn	Gly	Ile		Val	
347 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala 348 275 280 285 351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu									•	_		4		•		_		
348 275 280 285 351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu		Asn	Phe	Lys		Arq	His	Asn	Ile		Asp	Gly	Gly	Val			Ala	
351 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu				_		J					-	-	4					
		Asp	His	Tyr	Gln	Gln	Asn	Thr		Ile	Gly	Asp	Gly		Val	Leu	Leu	
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DATE: 06/27/2006

TIME: 10:43:38

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443 His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/583,591

VERIFICATION SUMMARY

DATE: 06/27/2006

PATENT APPLICATION: US/10/583,591 TIME: 10:43:39

Input Set : A:\PA0394 Sequence Listing.ST25.txt

Output Set: N:\CRF4\06272006\J583591.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number